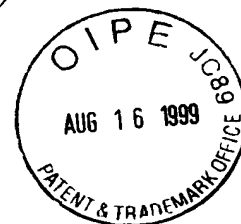SEQUENCE LISTING

<110> Gray, Joe W
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      Hwang, Soo In
      Godfrey, Tony
      Kowel, David
      Rommens, Johanna

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<141> 1997-07-15

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1

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<211> 723
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<223> Description of Artificial Sequence:1b11

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atttgattgg gtattgagca tgctctgaat gacagtcccc aaaaaggacc tcttatccgt  180
tcttcccctt ggggaagggc ttttgccact tccatgtcaa tgtggcagtt gagcttggaa  240
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aacatgtttt gggggaatgt tttgtttttc attttcttat tacctggcaa aataatccag  420
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acagtggacc tcaacgaagg agatgctgca cctgaaccca cwgaagcgaa actcaaaaga  540
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cttccagtat aggaaaccta agatacctag agcggctttt gggaacaatg ggctcatgcc  660
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<210> 3
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:cc49

<400> 3

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gaattggggc aagaagggag caccgacaac gacgattcga gttccgagaa ggagcttgga  1200
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3

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<211> 2605
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:cc43

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gatggcgatg agtttcgagt ggccgtggca gtatcgcttc ccacccttct ttacgttaca 180
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4

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<211> 1288
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:41.1

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<210> 6
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<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence:GCAP

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6

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<210> 7
<211> 1205
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:1b4

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8

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11

<213> Artificial Sequence

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tatccagaag ttttaatgat gcaccagaga ctggagcata aatacaatcc tgacgttcat 2220
aaaaactgtc gaaacaagtc cttgcttaga agtcgacgta ccggatgccc gccagcgttg 2280
ctgggaaaag atgtgcctcc cctctctagt ttctgtaaac ccaagcccaa gtctgctttc 2340
ccggcgcagt ccaaatccct gccatctgcg aaggggaagc agagccctcc tgggccaggc 2400
aaggcccctc tgacttcagg gatagactct agcactttag ccccaagtaa cctgaagtcc 2460

12

```
cacagaccac agcagaatgt gggggtccaa ggggccgcca ccaggcaaca gcaatctgag 2520
atgtttccta aaaccagtgt ttcccctgca ccggataaga caaaaagacc cgagacaaaa 2580
ttgaaacctc ttccagtagc tccttctcag cccacccctcg gcagcagtaa catcaatggt 2640
tccatcgact accccgccaa gaacgacagc ccgtgggcac ctccgggaag agactatttc 2700
tgtaatcgga gtgccagcaa tactgcagca gaatttggtg agccccttcc aaaaagactg 2760
aagtccagcg tggttgccct tgacgttgac cagcccgggg ccaattacag aagaggctat 2820
gaccttccca agtaccatat ggtcagaggc atcacatcac tgttaccgca ggactgtgtg 2880
tatccgtcgc aggcgctgcc tcccaaacca aggttcctga gctccagcga ggtcgattct 2940
ccaaatgtgc tgactgttca gaagccctat ggtggctccg ggccacttta cacttgtgtg 3000
cctgctggta gtccagcatc cagctcgacg ttagaaggtc ttggtggatg tcagtgctta 3060
ctccccatga aattaaattt tacttcatcc tttgagaagc gaatggtgaa agctactgaa 3120
ataagctgtg attgtactgt acataaaaca tatgaggaat ctgcaaggaa cactacagtt 3180
gtgtaa                                                             3186
```

<210> 11
<211> 1061
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:ZABC1 Protein

<400> 11

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Met Gln Ser Lys Val Thr Gly Asn Met Pro Thr Gln Ser Leu Leu Met
  1               5                  10                  15

Tyr Met Asp Gly Pro Glu Val Ile Gly Ser Ser Leu Gly Ser Pro Met
                20                  25                  30

Glu Met Glu Asp Ala Leu Ser Met Lys Gly Thr Ala Val Val Pro Phe
            35                  40                  45

Arg Ala Thr Gln Glu Lys Asn Val Ile Gln Ile Glu Gly Tyr Met Pro
        50                  55                  60

Leu Asp Cys Met Phe Cys Ser Gln Thr Phe Thr His Ser Glu Asp Leu
 65                  70                  75                  80

Asn Lys His Val Leu Met Gln His Arg Pro Thr Leu Cys Glu Pro Ala
                85                  90                  95

Val Leu Arg Val Glu Ala Glu Tyr Leu Ser Pro Leu Asp Lys Ser Gln
            100                 105                 110

Val Arg Thr Glu Pro Pro Lys Glu Lys Asn Cys Lys Glu Asn Glu Phe
        115                 120                 125

Ser Cys Glu Val Cys Gly Gln Thr Phe Arg Val Ala Phe Asp Val Glu
```

13

Ile His Met Arg Thr His Lys Asp Ser Phe Thr Tyr Gly Cys Asn Met
145   150   155   160

Cys Gly Arg Xaa Xaa Xaa Xaa Pro Trp Phe Leu Lys Asn His Met Arg
                    165   170   175

Thr His Asn Gly Lys Ser Gly Ala Arg Ser Lys Leu Gln Gln Gly Leu
                    180   185   190

Glu Ser Ser Pro Ala Thr Ile Asn Glu Val Val Gln Val His Ala Ala
                    195   200   205

Glu Ser Ile Ser Ser Pro Tyr Lys Ile Cys Met Val Cys Gly Phe Leu
          210   215   220

Phe Pro Asn Lys Glu Ser Leu Ile Glu His Arg Lys Val His Thr Lys
225   230   235   240

Lys Thr Ala Phe Gly Thr Ser Ser Ala Gln Thr Asp Ser Pro Gln Gly
                    245   250   255

Gly Met Pro Ser Ser Arg Glu Asp Phe Leu Gln Leu Phe Asn Leu Arg
                    260   265   270

Pro Lys Ser His Pro Glu Thr Gly Lys Lys Pro Val Arg Cys Ile Pro
          275   280   285

Gln Leu Asp Pro Phe Thr Thr Phe Gln Ala Trp Gln Leu Ala Thr Lys
          290   295   300

Gly Lys Val Ala Ile Cys Gln Glu Val Lys Glu Ser Gly Gln Glu Gly
305   310   315   320

Ser Thr Asp Asn Asp Asp Ser Ser Ser Glu Lys Glu Leu Gly Glu Thr
                    325   330   335

Asn Lys Gly Ser Cys Ala Gly Leu Ser Gln Glu Lys Glu Lys Cys Lys
                    340   345   350

His Ser His Gly Glu Ala Pro Ser Val Asp Ala Asp Pro Lys Leu Pro
          355   360   365

Ser Ser Lys Glu Lys Pro Thr His Cys Ser Glu Cys Gly Lys Ala Phe
          370   375   380

Arg Thr Tyr His Gln Leu Val Leu His Ser Arg Val His Lys Lys Asp

14

Arg Arg Ala Gly Ala Glu Ser Pro Thr Met Ser Val Asp Gly Arg Gln
            405   410   415

Pro Gly Thr Cys Ser Pro Asp Leu Ala Ala Pro Leu Asp Glu Asn Gly
            420   425   430

Ala Val Asp Arg Gly Glu Gly Gly Ser Glu Asp Gly Ser Glu Asp Gly
            435   440   445

Leu Pro Glu Gly Ile His Leu Asp Lys Asn Asp Asp Gly Gly Lys Ile
    450   455   460

Lys His Leu Thr Ser Ser Arg Glu Cys Ser Tyr Cys Gly Lys Phe Phe
465   470   475   480

Arg Ser Asn Tyr Tyr Leu Asn Ile His Leu Arg Thr His Thr Gly Glu
            485   490   495

Lys Pro Tyr Lys Cys Glu Phe Cys Glu Tyr Ala Ala Ala Gln Lys Thr
            500   505   510

Ser Leu Arg Tyr His Leu Glu Arg His His Lys Glu Lys Gln Thr Asp
            515   520   525

Val Ala Ala Glu Val Lys Asn Asp Gly Lys Asn Gln Asp Thr Glu Asp
    530   535   540

Ala Leu Leu Thr Ala Asp Ser Ala Gln Thr Lys Asn Leu Lys Arg Phe
545   550   555   560

Phe Asp Gly Ala Lys Asp Val Thr Gly Ser Pro Pro Ala Lys Gln Leu
            565   570   575

Lys Glu Met Pro Ser Val Phe Gln Asn Val Leu Gly Ser Ala Val Leu
            580   585   590

Ser Pro Ala His Lys Asp Thr Gln Asp Phe His Lys Asn Ala Ala Asp
            595   600   605

Asp Ser Ala Asp Lys Val Asn Lys Asn Pro Thr Pro Ala Tyr Leu Asp
    610   615   620

Leu Leu Lys Lys Arg Ser Ala Val Glu Thr Gln Ala Asn Asn Leu Ile
625   630   635   640

Cys Arg Thr Lys Ala Asp Val Thr Pro Pro Pro Asp Gly Ser Thr Thr

15

```
His Asn Leu Glu Val Ser Pro Lys Glu Lys Gln Thr Glu Thr Ala Ala
            660             665             670

Asp Cys Arg Tyr Arg Pro Ser Val Asp Cys His Glu Lys Pro Leu Asn
            675             680             685

Leu Ser Val Gly Ala Leu His Asn Cys Pro Ala Ile Ser Leu Ser Lys
            690             695             700

Ser Leu Ile Pro Ser Ile Thr Cys Pro Phe Cys Thr Phe Lys Thr Phe
705             710             715             720

Tyr Pro Glu Val Leu Met Met His Gln Arg Leu Glu His Lys Tyr Asn
            725             730             735

Pro Asp Val His Lys Asn Cys Arg Asn Lys Ser Leu Leu Arg Ser Arg
            740             745             750

Arg Thr Gly Cys Pro Pro Ala Leu Leu Gly Lys Asp Val Pro Pro Leu
            755             760             765

Ser Ser Phe Cys Lys Pro Lys Pro Lys Ser Ala Phe Pro Ala Gln Ser
    770             775             780

Lys Ser Leu Pro Ser Ala Lys Gly Lys Gln Ser Pro Pro Gly Pro Gly
785             790             795             800

Lys Ala Pro Leu Thr Ser Gly Ile Asp Ser Ser Thr Leu Ala Pro Ser
            805             810             815

Asn Leu Lys Ser His Arg Pro Gln Gln Asn Val Gly Val Gln Gly Ala
            820             825             830

Ala Thr Arg Gln Gln Gln Ser Glu Met Phe Pro Lys Thr Ser Val Ser
            835             840             845

Pro Ala Pro Asp Lys Thr Lys Arg Pro Glu Thr Lys Leu Lys Pro Leu
    850             855             860

Pro Val Ala Pro Ser Gln Pro Thr Leu Gly Ser Ser Asn Ile Asn Gly
865             870             875             880

Ser Ile Asp Tyr Pro Ala Lys Asn Asp Ser Pro Trp Ala Pro Pro Gly
            885             890             895

Arg Asp Tyr Phe Cys Asn Arg Ser Ala Ser Asn Thr Ala Ala Glu Phe
```

16

```
                  900                      905                      910

        Gly Glu Pro Leu Pro Lys Arg Leu Lys Ser Ser Val Val Ala Leu Asp
                915                 920                 925

        Val Asp Gln Pro Gly Ala Asn Tyr Arg Arg Gly Tyr Asp Leu Pro Lys
                930                 935                 940

        Tyr His Met Val Arg Gly Ile Thr Ser Leu Leu Pro Gln Asp Cys Val
        945                 950                 955                 960

        Tyr Pro Ser Gln Ala Leu Pro Pro Lys Pro Arg Phe Leu Ser Ser Ser
                        965                 970                 975

        Glu Val Asp Ser Pro Asn Val Leu Thr Val Gln Lys Pro Tyr Gly Gly
                980                 985                 990

        Ser Gly Pro Leu Tyr Thr Cys Val Pro Ala Gly Ser Pro Ala Ser Ser
                995                 1000                1005

        Ser Thr Leu Glu Gly Leu Gly Gly Cys Gln Cys Leu Leu Pro Met Lys
            1010                1015                1020

        Leu Asn Phe Thr Ser Ser Phe Glu Lys Arg Met Val Lys Ala Thr Glu
        1025                1030                1035                1040

        Ile Ser Cys Asp Cys Thr Val His Lys Thr Tyr Glu Glu Ser Ala Arg
                        1045                1050                1055

        Asn Thr Thr Val Val
                    1060


        <210> 12
        <211> 3066
        <212> DNA
        <213> Artificial Sequence

        <220>
        <223> Description of Artificial Sequence:1b1

        <400> 12
        ggaaacagct atgaccatga ttacgccaag ctcgaaatta accctcacta aagggaacaa 60
        aagctggagc tccaccgcgg tggcggccgc tctagaacta gtggatcccc cgggctgcag 120
        gaattcggca cgaggctcca ccgacagcca ggcactgggc agcacgcact ggagacccag 180
        gaccctgtgc aggagcagct ccgggtgaca cgaggggact gaagatactc ccacaggggc 240
        tcagcaggag caatgggtaa ccaaatgagt gttccccaaa gagttgaaga ccaagagaat 300
        gaaccagaag cagagactta ccaggacaac gcgtctgctc tgaacggggt tccagtggtg 360
```

17

```
gtgtcgaccc acacagttca gcacttagag gaagtcgact tgggaataag tgtcaagacg 420
gataatgtgg ccacttcttc ccccgagaca acggagataa gtgctgttgc ggatgccaac 480
ggaaagaatc ttgggaaaga ggccaaaccc gaggcaccag ctgctaaatc tcgttttttc 540
ttgatgctct ctcggcctgt accaggacgt accggagacc aagccgcaga ttcatcccta 600
ggatcagtga agcttgatgt cagctccaat aaagctccag cgaacaaaga cccaagtgag 660
agctggacac ttccggtggc agctggaccg gggcaggaca cagataaaac cccagggcac 720
gccccggccc aagacaaggt cctctctgcc gccagggatc ccacgcttct cccacctgag 780
acaggggag caggaggaga agctccctcc aagcccaagg actccagctt ttttgacaaa 840
ttcttcaagc tggacaaggg acaggaaaag gtgccaggtg acagccaaca ggaagccaag 900
agggcagagc atcaagacaa ggtggatgag gttcctggct tatcagggca gtccgatgat 960
gtccctgcag ggaaggacat agttgacggc aaggaaaaag aaggacaaga acttggaact 1020
gcggattgct ctgtccctgg ggacccagaa ggactggaga ctgcaaagga cgattcccag 1080
gcagcagcta tagcagagaa taataattcc atcatgagtt tctttaaaac tctggtttca 1140
cctaacaaag ctgaaacaaa aaaggaccca gaagacacgg gtgctgaaaa gtcacccacc 1200
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cctgcgaagg aaggcaccaa ggagaaatca ggacccacct ctctgcctct gggcaaactg 1380
tttttggaaaa agtcagttaa agaggactca gtccccacag gtgcggagga gaatgtggtg 1440
tgtgagtcac cagtagagat tataaagtcc aaggaagtag aatcagcctt acaaacagtg 1500
gacctcaacg aaggagatgc tgcacctgaa cccacagaag cgaaactcaa aagagaagaa 1560
agcaaaccaa gaacctctct gatggcgttt ctcagacaaa tgtcagtgaa aggggatgga 1620
gggatcaccc actcagaaga aataaatggg aaagactcca gctgccaaac atcagactcc 1680
acagaaaaga ctatcacacc gccagagcct gaaccaacag gagcaccaca gaagggtaaa 1740
gagggctcct cgaaggacaa gaagtcagca gccgagatga acaagcagaa gagcaacaag 1800
caggaagcca aagaaccagc ccagtgcaca gagcaggcca cggtggacac gaactcactg 1860
cagaatgggg acaagctcca aaagagacct gagaagcggc agcagtccct tgggggcttc 1920
tttaaaggcc tgggaccaaa gcggatgttg gatgctcaag tgcaaacaga cccagtatcc 1980
atcggaccag ttggcaaacc caagtaaaca aatcagcacg gttcccacca ggttctcctg 2040
ccaccaagat gtgttctcct tactccatct cctccccaaa cacgctccat gtatatattc 2100
ttctgatggc cagcaaatga aattctgcct agaaattaag cccgagctgt tgtatattga 2160
ggtgtattat ttacgtctct ggtccagtct tttctggcaa ataacagtaa agatggttta 2220
gcaggtcacc tagttgggtc agaagagtcg atgatcacca agcaggaaag ggagggaata 2280
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gggagacgaa aatccatttg tttaaattca catctcaagg agggagaacc cgggctgtgt 2760
tgggtggttg ccaatttcct agaacggaat gtgtggggta tagaaaaagg aatgaataag 2820
cgttgttttt caaatagggt ccttgtaagt tattgatgag agggaaaaga ttgactgggg 2880
agggcttaaa atgatttggg aaaacaattg cttttgaggc tcagtgacaa cggcaaagat 2940
tacaacttaa aaaaaaaaaa aaaaaaactc gagactagtt ctctctctct ctcgtgccga 3000
attcgatatc aagcttatcg ataccgtcga cctcgagggg gggcccggta cccaattcgc 3060
cctata                                                              3066
```

<210> 13
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<211> 939
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Genomic
      Sequence from BAC Clone 97 Filtered Query Sequence

<400> 13
tgtgatattg attcatgccc tcttgcacct tgccaaacat cacacgcttg ccatccagtc 60
cactcgattt tggcagtgca gatgaaaaac tgggaaccat ttgtgttgag tccagcaaga 120
tgccaggacc tgcatgtttc agaacgaagt tcttcatcat ccaatttctc cctgtatatg 180
ggcttaccac nactgccgtt aagtcgtgtn aagtcaccac tcaggtacat aatggaataa 240
ttctgcaaag gcaggagnca ctttctctcc agtgctcaga ccatgaaagt tttctgatgt 300
ctttggaact ttgtctgcaa atagctcgaa ggagacatgg cctaaaggct cgccatctgc 360
ggtgatattg naacatggta gggctgaccg tggctgtggc catgactttt tagantnnnn 420
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 480
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnncccaat gcgggacaga gaatcnaaga 540
aactgtatta gggaaagggt cctgagttta tgccaaagtt tcccagattg gtttccattg 600
aaacgtagct ctgtgagata ccatcaggtg ttatgtgaag aaatgtctgt gtagtcaaat 660
atgtttgagt gagtgagcct gagctgagca agactttact gcaagacttc ccatcttctg 720
tcccttttta tgctaatggg taacacaaac tccaaaagtg gggtgtacag catgaggcat 780
taacaaaaat ttattggacc ccacacacnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 840
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 900
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnctctc                        939

<210> 14
<211> 112
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Subject Seq -
      Rat Cyclophillin 64-175

<400> 14
ttcgacatca cggctgatgg cgagcccttg ggtcgcgtct gcttcgagct gtttgcagac 60
aaagttccaa agacagcaga aaactttcgt gctctgagca ctggggagaa ag          112

<210> 15
<211> 106
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:e:Subject Seq -
      Rat Cyclophilli404-348

.

```
<400> 15
tgctggacca aacacaaatg gttcccagtt ttttatctgc actgccaaga ctgagtgggg 60
ctggatggca agcatgtggt ctttgggaag gtgaaagaag gcatga                106


<210> 16
<211> 38
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence:e:Subject Seq -
     Rat Cyclophillin 299-336


<400> 16
agaacttcat cctgaagcat acaggtcctg gcatcttg                         38


<210> 17
<211> 28
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence:e:Subject Seq -
     Rat Cyclophillin 193-220


<400> 17
tcctcctttc acagaattat tccaggat                                    28


<210> 18
<211> 112
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence:Query Seq ID NO
     13 261-372


<400> 18
tncaatatca ccgcagatgg cgagccttta ggccatgtct ccttcgagct atttgcagac 60
aaagttccaa agacatcaga aaactttcat ggtctgagca ctggagagaa ag         112


<210> 19
<211> 106
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence:Query Seq ID NO
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13 13-117

<400> 19
tgctggactc aacacaaatg gttcccagtt tttcatctgc actgccaaaa tcgagtggga 60
ctggatggca agcgtgtgat gtttggcaag gtgcaagagg gcatga   106

<210> 20
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Query Seq ID NO
      13116-153

<400> 20
agaacttcgt tctgaaacat gcaggtcctg gcatcttg   38

<210> 21
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Query Seq ID NO
      13 229-256

<400> 21
tcctgccttt gcagaattat tccattat   28

<210> 22
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Forward primer

<400> 22
ttggcattgg tatcaggtag ctg   23

<210> 23
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward

Primer

<400> 23
ttggagcaga gaggggattg tgtg   24

<210> 24
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:: Forward
      primer

<400> 24
aatcccctca aaccctgctg ctac   24

<210> 25
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward
      Primer

<400> 25
tggagcctga acttctgcaa tc   22

<210> 26
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:: Forward
      primer

<400> 26
ccgggatacc gacattg   17

<210> 27
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward

22

Primer

<400> 27
tgcacataaa acagccagc

19

<210> 28
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:: Forward
      primer

<400> 28
ttggaatcaa tggagcaaaa

20

<210> 29
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward
      Primer

<400> 29
agctttaccc aatgtggtcc

20

<210> 30
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:: Forward
      primer

<400> 30
gtggtgaaca ccaataaatg g

21

<210> 31
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward

23

```
          Primer

<400> 31
aagcaaataa aaccaataaa ctcg                                        24


<210> 32
<211> 21
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence:: Forward
      primer

<400> 32
caagatctga ccccgtcaat c                                           21


<210> 33
<211> 25
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence: Backward
      Primer

<400> 33
gacttcttca ggaaagagat cagtg                                       25


<210> 34
<211> 23
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence:: Forward
      primer


<400> 34
gccatgtacc cacctgaaaa atc                                         23


<210> 35
<211> 24
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence: Backward
```

Primer

<400> 35
tcagaacacc cgtgcagaat taag   24

<210> 36
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:: Forward
      primer

<400> 36
cctaaaactt ggtgcttaaa tcta   24

<210> 37
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward
      Primer

<400> 37
gtctcacaag gcagatgtgg   20

<210> 38
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:: Forward
      primer

<400> 38
tttgtgtatg ttgagccatc   20

<210> 39
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward

Primer

<400> 39
cttccaatct cattctatga gg   22

<210> 40
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:Forward Primer

<400> 40
gcttgtttaa gtgtcactag gg   22

<210> 41
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward
      Primer

<400> 41
cactctggta aatgaccttt gtc   23

<210> 42
<211> 21   .
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:: Forward
      primer

<400> 42
cctacaccat tccaactttg g   21

<210> 43
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward
      Primer

```
<400> 43
gccagatgta tgtttgctac ggaac                                    25


<210> 44
<211> 22
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence:: Forward
      primer


<400> 44
tctcaaacct gtccacttct tg                                       22


<210> 45
<211> 19
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence: Backward
      Primer


<400> 45
ctgctgtggt ggagaatgg                                           19


<210> 46
<211> 24
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence:: Forward
      primer


<400> 46
tgtcctcctt ctccctcatc ctac                                     24


<210> 47
<211> 22
<212> DNA
<213> Artificial Sequence


<220>
<223> Description of Artificial Sequence: Backward
      Primer
```

<400> 47
aatgcctcca ctcacaggaa tg   22

<210> 48
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:: Forward
     primer

<400> 48
cctcttcagt gtcttcctat tga   23

<210> 49
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Backward
     Primer

<400> 49
gggaggaggt tgtaggcaac   20

<210> 50
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Forward primer

<400> 50
agcaaagcaa aggtggcaca c   21

<210> 51
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     Primer

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24

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20

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